

THE SPECIFICATION:

Please replace the paragraph beginning at page 15, line 3 with the following paragraph:

--**Figure 1** is a representation of the sequence, exon composition and pre-mRNA structure of the alternative 5'UTRs of mouse *GLII*. *Panel A*, sequence alignment of the three alternative *GLII* 5'UTR variants (denoted α -UTR (SEQ ID NO: 61), β -UTR (SEQ ID NO: 62) and γ -UTR (SEQ ID NO: 63)) expressed in mouse. The novel 119 bp sequence of exon 1a is shown in bold lowercase lettering. The ATG codons denoting the beginning of uORFs are underlined and the main ORF encoding *GLII* is shown bold uppercase lettering. The intron/exon boundaries are indicated by arrows. *Panel B*, schematic showing the exon composition of the alternative 5'UTRs and the organization of the pre-mRNA from which they are derived. Exons are denoted by open boxes and introns by solid lines with intron size shown. The translation start site (ATG) of the main ORF is located in exon 2 and indicated by a bent arrow.--

Please replace the paragraph beginning at page 15, line 28, with the following paragraph:

--**Figure 4** is a representation of the sequence, exon composition and pre-mRNA structure of alternative human *GLII* 5'UTRs, *Panel A*, sequence alignment of the alternative *GLII* 5'UTR variants identified in human tissues (denoted β -UTR (SEQ ID NO: 64) and γ -UTR (SEQ ID NO: 65)). The novel 144 bp sequence of exon 1a is shown in bold lowercase lettering. The ATG codons denoting the beginning of uORFs are underlined and the main ORF encoding *GLII* is shown in bold uppercase lettering. The intron/exon boundaries are indicated by arrows. *Panel B*, schematic showing the exon composition of the alternative 5'UTRs and the organization of the pre-mRNA from which they are derived. Exons are denoted by open boxes and introns by solid lines with intron size shown. The translation start site (ATG) of the main ORF is located in